

SEQUENCE LISTING

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<120> RECOMBINANT FUSOBACTERIUM NECROPHORUM LEUKOTOXIN VACCINE AND PREPARATION THEREOF

<130> PCT-30296

<160> 15

<170> PatentIn version 3.0

<210> 1

<211> 3241

<212> PRT

<213> *Fusobacterium necrophorum*

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Lys Ile Glu Lys Lys Asp Asn Val Tyr Asp Ile Thr Thr Asn Lys Ile
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Gln Gly Glu Asn Ala Phe Asn Ser Phe Asn Arg Phe Ala Leu Thr Glu
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Asn Asn Ile Ala Asn Leu Tyr Phe Gly Glu Lys Asn Ser Thr Gly Val
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Asn Asn Leu Phe Asn Phe Val Asn Gly Lys Ile Glu Val Asp Gly Ile
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PCT302~1

Ile Asn Gly Ile Arg Glu Asn Lys Ile Gly Gly Asn Leu Tyr Phe Leu
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Ser Ser Glu Gly Met Ala Val Gly Lys Asn Gly Val Ile Asn Ala Gly
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Ser Phe His Ser Ile Ile Pro Lys Gln Asp Asp Phe Lys Lys Ala Leu
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Glu Glu Ala Lys His Gly Lys Val Phe Asn Gly Ile Ile Pro Val Asp
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Gly Lys Val Lys Ile Pro Leu Asn Pro Asn Gly Ser Ile Thr Val Glu
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Gly Lys Ile Asn Ala Val Glu Gly Ile Gly Leu Tyr Ala Ala Asp Ile
195 200 205

Arg Leu Lys Asp Thr Ala Ile Leu Lys Thr Gly Ile Thr Asp Phe Lys
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Asn Leu Val Asn Ile Ser Asp Arg Ile Asn Ser Gly Leu Thr Gly Asp
225 230 235 240

Leu Lys Ala Thr Lys Thr Lys Ser Gly Asp Ile Ile Leu Ser Ala His
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Ile Asp Ser Pro Gln Lys Ala Met Gly Lys Asn Ser Thr Val Gly Lys
260 265 270

Arg Ile Glu Glu Tyr Val Lys Gly Asn Thr Lys Ala Asn Ile Glu Ser
275 280 285

Asp Ala Val Leu Glu Ala Asp Gly Asn Ile Lys Ile Ser Ala Lys Ala
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Thr Asn Gly Arg Phe Ile Lys Lys Glu Gly Glu Lys Glu Thr Tyr Asn
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Thr Pro Leu Ser Leu Ser Asp Val Glu Ala Ser Val Arg Val Asn Lys
325 330 335

Gly Lys Val Ile Gly Lys Asn Val Asp Ile Thr Ala Glu Ala Lys Asn
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Phe Tyr Asp Ala Thr Leu Val Thr Lys Leu Ala Lys His Ser Phe Ser
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Phe Val Thr Gly Ser Ile Ser Pro Ile Asn Leu Asn Gly Phe Leu Gly
370 375 380

Leu Leu Thr Ser Lys Ser Ser Val Val Ile Gly Lys Asp Ala Lys Val
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Glu Ala Thr Glu Gly Lys Ala Asn Ile His Ser Tyr Ser Gly Val Arg
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Ala Thr Met Gly Ala Ala Thr Ser Pro Leu Lys Ile Thr Asn Leu Tyr
420 425 430

Leu Glu Lys Ala Asn Gly Lys Leu Leu Ser Ile Gly Ala Gly Tyr Ile
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Ser Ala Lys Ser Asn Ser Asn Val Thr Ile Glu Gly Glu Val Lys Ser
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Lys Gly Arg Ala Asp Ile Thr Ser Lys Ser Glu Asn Thr Ile Asp Ala

PCT302~1

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Gly Ala Lys Val Glu Ser Glu Thr Asp Asp Val Asn Val Arg Ser Glu			
515 520 525			
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Gly Asn Gly Val Val Ala Ala Asn Ile Ser Asn Tyr Asn Ala Ser Ser			
545 550 555 560			
Arg Ile Asp Val Asp Gly Tyr Leu His Ala Lys Lys Arg Leu Asn Val			
565 570 575			
Glu Ala His Asn Ile Thr Lys Asn Ser Val Leu Gln Thr Gly Ser Asp			
580 585 590			
Leu Gly Thr Ser Lys Phe Met Asn Asp His Val Tyr Glu Ser Gly His			
595 600 605			
Leu Lys Ser Ile Leu Asp Ala Ile Lys Gln Arg Phe Gly Gly Asp Ser			
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Val Asn Glu Glu Ile Lys Asn Lys Leu Thr Asn Leu Phe Ser Val Gly			
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Val Ser Ala Thr Ile Ala Asn His Asn Asn Ser Ala Ser Val Ala Ile			
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Gly Glu Ser Gly Arg Leu Ser Ser Gly Val Glu Gly Ser Asn Val Arg			
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Ala Leu Asn Glu Ala Gln Asn Leu Arg Ala Thr Thr Ser Ser Gly Ser			
675 680 685			
Val Ala Val Arg Lys Glu Glu Lys Lys Leu Ile Gly Asn Ala Ala			
690 695 700			
Val Phe Tyr Gly Asn Tyr Lys Asn Asn Ala Ser Val Thr Ile Ala Asp			
705 710 720			
His Ala Glu Leu Val Ser Glu Gly Lys Ile Asp Ile Asn Ser Glu Asn			
725 730 735			
Lys Ile Glu Tyr Lys Asn Pro Ser Lys Met Ala Lys Ser Val Ile Asp			
740 745 750			
Lys Leu Glu Leu Leu Lys Arg Ala Phe Gly Lys Glu Thr Lys Thr Pro			
755 760 765			
Glu Tyr Asp Pro Lys Asp Ile Glu Ser Ile Glu Lys Leu Leu Asn Ala			
770 775 780			
Phe Ser Glu Lys Leu Asp Gly Lys Pro Glu Leu Leu Leu Asn Gly Glu			
785 790 795 800			
Arg Met Thr Ile Ile Leu Pro Asp Gly Thr Ser Lys Thr Gly Thr Ala			
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Ile Glu Ile Ala Asn Tyr Val Gln Gly Glu Met Lys Lys Leu Glu Glu			
820 825 830			

PCT302~1

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 835 840 845

Ile Lys Glu Thr Leu Asn Phe Thr Gly Val Gly Asn Tyr Ala Asn Phe
 850 855 860

His Thr Phe Thr Ser Ser Gly Ala Asn Gly Glu Arg Asp Val Ser Ser
 865 870 875 880

Val Gly Gly Ala Val Ser Trp Val Glu Gln Glu Asn Tyr Ser Lys Val
 885 890 895

Ser Val Gly Lys Gly Ala Lys Leu Ala Ala Lys Lys Asp Leu Asn Ile
 900 905 910

Lys Ala Ile Asn Lys Ala Glu Thr Val Asn Leu Val Gly Asn Ile Gly
 915 920 925

Leu Ala Arg Ser Ser Thr Ser Gly Ser Ala Val Gly Gly Arg Leu Asn
 930 935 940

Val Gln Arg Ser Lys Asn Ser Ala Ile Val Glu Ala Lys Glu Lys Ala
 945 950 955 960

Glu Leu Ser Gly Glu Asn Ile Asn Ala Asp Ala Leu Asn Arg Leu Phe
 965 970 975

His Val Ala Gly Ser Phe Asn Gly Gly Ser Gly Gly Asn Ala Ile Asn
 980 985 990

Gly Met Gly Ser Tyr Ser Gly Gly Ile Ser Lys Ala Arg Val Ser Ile
 995 1000 1005

Asp Asp Glu Ala Tyr Leu Lys Ala Asn Lys Ile Ala Leu Asn
 1010 1015 1020

Ser Lys Asn Asp Thr Ser Val Trp Asn Ala Ala Gly Ser Ala Gly
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Ile Gly Thr Lys Asn Ala Ala Val Gly Val Ala Val Ala Val Asn
 1040 1045 1050

Asp Tyr Asp Ile Ser Asn Lys Ala Ser Ile Glu Asp Asn Asp Glu
 1055 1060 1065

Gly Gln Ser Lys Tyr Asp Lys Asn Lys Asp Asp Glu Val Thr Val
 1070 1075 1080

Thr Ala Glu Ser Leu Glu Val Asp Ala Lys Thr Thr Gly Thr Ile
 1085 1090 1095

Asn Ser Ile Ser Val Ala Gly Gly Ile Asn Lys Val Gly Ser Lys
 1100 1105 1110

Pro Ser Glu Glu Lys Pro Lys Ser Glu Glu Arg Pro Glu Gly Phe
 1115 1120 1125

Phe Gly Lys Ile Gly Asn Lys Val Asp Ser Val Lys Asn Lys Ile
 1130 1135 1140

Thr Asp Ser Met Asp Ser Leu Thr Glu Lys Ile Thr Asn Tyr Ile
 1145 1150 1155

Ser Glu Gly Val Lys Lys Ala Gly Asn Leu Pro Ser Asn Val Ser
 1160 1165 1170

His Thr Pro Asp Lys Gly Pro Ser Phe Ser Leu Gly Ala Ser Gly
 1175 1180 1185

PCT302~1

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 Val Thr Ser Ser Asp Ser Thr 1220 Phe Val Gly Ala Trp 1225 Gly Gly Ser 1230
 Ala Ala Leu Gln Trp Asn His 1235 Ile Gly Ser Gly Asn 1240 Ser Asn Ile 1245
 Ser Ala Gly Leu Ala Gly Ala 1250 Ala Ala Val Asn Asn 1255 Ile Gln Ser 1260
 Lys Thr Ser Ala Leu Val Lys 1265 Asn Ser Asp Ile Arg 1270 1275 Asn Ala Asn
 Lys Phe Lys Val Asn Ala Leu 1280 Ser Gly Gly Thr Gln 1285 Val Ala Ala 1290
 Gly Ala Gly Leu Glu Ala Val 1295 Lys Glu Ser Gly Gly 1300 Gln Gly Lys 1305
 Ser Tyr Leu Leu Gly Thr Ser 1310 Ala Ser Ile Asn Leu 1315 Val Asn Asn 1320
 Glu Val Ser Ala Lys Ser Glu 1325 Asn Asn Thr Val Ala 1330 Gly Glu Ser 1335
 Glu Ser Gln Lys Met Asp Val 1340 Asp Val Thr Ala Tyr 1345 Gln Ala Asp 1350
 Thr Gln Val Thr Gly Ala Leu 1355 Asn Leu Gln Ala Gly 1360 Lys Ser Asn 1365
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 Ala Asp Ala Lys Ala Leu Leu 1400 Ala Thr Thr Gln Val 1405 1410 Thr Ala Ala
 Val Thr Thr Gly Gly Thr Ile 1415 Ser Ser Gly Ala Gly 1420 1425 Leu Gly Asn
 Tyr Gln Gly Ala Val Ser Val 1430 Asn Lys Ile Asp Asn 1435 1440 Asp Val Glu
 Ala Ser Val Asp Lys Ser Ser 1445 Ile Glu Gly Ala Asn 1450 1455 Glu Ile Asn
 Val Ile Ala Lys Asp Val Lys 1460 Gly Ser Ser Asp Leu 1465 Ala Lys Glu 1470
 Tyr Gln Ala Leu Leu Asn Gly 1475 Lys Asp Lys Lys Tyr 1480 1485 Leu Glu Asp
 Arg Gly Ile Asn Thr Thr Gly 1490 Asn Gly Tyr Tyr Thr 1495 1500 Lys Glu Gln
 Leu Glu Lys Ala Lys Lys Lys 1505 Glu Gly Ala Val Ile 1510 1515 Val Asn Ala
 Ala Leu Ser Val Ala Gly Thr Asp Lys Ser Ala Gly Gly Val Ala

PCT302~1

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Gly Ser Asn Lys Glu Ala	Gly Glu Asp Lys Ile His	Ala Lys His
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Val Asn Val Glu Ala Lys	Ser Thr Val Val	Asn Ala Ala
1565	1570	1575
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Gly Ala Trp Gln Asp Leu	Ser Asp Thr Ile Ala	Lys Val Asp
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Lys Gly Arg Ile Ser Ala	Ser Leu Asn Val Asn	Ala Asn Asn
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Ser Ile Leu Gly Val Asn	Val Ala Gly Thr Ile Ala	Gly Ser Leu
1625	1630	1635
Ser Thr Ala Val Gly Ala	Phe Ala Asn Asn Thr	Leu His Asn
1640	1645	1650
Lys Thr Ser Ala Leu Ile	Thr Gly Thr Lys Val Asn	Pro Phe Ser
1655	1660	1665
Gly Lys Asn Thr Lys Val	Asn Val Gln Ala Leu Asn	Asp Ser His
1670	1675	1680
Ile Thr Asn Val Ser Ala	Gly Ala Ala Ser Ile	Lys Gln Ala
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Gly Ile Gly Gly Met Val	Ser Val Asn Arg Gly	Ser Glu Thr
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Glu Ala Leu Val Ser Asp	Glu Phe Glu Gly Val	Ser Ser Phe
1715	1720	1725
Asn Val Asp Ala Lys Asp	Gln Lys Thr Ile Asn Thr	Ile Ala Gly
1730	1735	1740
Asn Ala Asn Gly Gly Lys	Ala Gly Val Gly Ala	Thr Val Ala
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His Thr Asn Ile Gly Lys	Ser Val Ile Ala Ile	Val Lys Asn
1760	1765	1770
Ser Lys Ile Thr Thr Ala	Asp Gln Asp Arg Lys	Asn Ile Asn
1775	1780	1785
Val Thr Ala Lys Asp Tyr	Thr Met Thr Asn Thr	Ile Ala Val Gly
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Val Gly Gly Ala Lys Gly	Ala Ser Val Gln Gly	Ala Ser Ala
1805	1810	1815
Thr Thr Leu Asn Lys Thr	Val Ser Ser His Val	Asp Gln Thr Asp
1820	1825	1830
Ile Asp Lys Asp Leu Glu	Glu Asn Asn Gly Asn	Lys Glu Lys
1835	1840	1845
Ala Asn Val Asn Val Leu	Glu Asn Thr Ser Gln	Val Val Thr
1850	1855	1860

PCT302~1

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 Ala Gly Val Ala Val Asn Lys Ile Thr Gln Asn Thr Ser Ala His
 1880 1885 1890
 Ile Lys Asn Ser Thr Gln Asn Val Arg Asn Ala Leu Val Lys Ser
 1895 1900 1905
 Lys Ser His Ser Ser Ile Lys Thr Ile Gly Ile Gly Ala Gly Val
 1910 1915 1920
 Gly Ala Gly Gly Ala Gly Val Thr Gly Ser Val Ala Val Asn Lys
 1925 1930 1935
 Ile Val Asn Asn Thr Ile Ala Glu Leu Asn His Ala Lys Ile Thr
 1940 1945 1950
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 1955 1960 1965
 Ala Asn Tyr Ala Gly Thr Val Ser Gly Val Ala Arg Ala Ala Ile
 1970 1975 1980
 Gly Ala Ser Thr Ser Val Asn Glu Ile Thr Gly Ser Thr Lys Ala
 1985 1990 1995
 Tyr Val Lys Asp Ser Thr Val Ile Ala Lys Glu Glu Thr Asp Asp
 2000 2005 2010
 Tyr Ile Thr Thr Gln Gly Gln Val Asp Lys Val Val Asp Lys Val
 2015 2020 2025
 Phe Lys Asn Leu Asn Ile Asn Glu Asp Leu Ser Gln Lys Arg Lys
 2030 2035 2040
 Ile Ser Asn Lys Lys Gly Phe Val Thr Asn Ser Ser Ala Thr His
 2045 2050 2055
 Thr Leu Lys Ser Leu Leu Ala Asn Ala Ala Gly Ser Gly Gln Ala
 2060 2065 2070
 Gly Val Ala Gly Thr Val Asn Ile Asn Lys Val Tyr Gly Glu Thr
 2075 2080 2085
 Glu Ala Leu Val Glu Asn Ser Ile Leu Asn Ala Lys His Tyr Ser
 2090 2095 2100
 Val Lys Ser Gly Asp Tyr Thr Asn Ser Ile Gly Val Val Gly Ser
 2105 2110 2115
 Val Gly Val Gly Gly Asn Val Gly Val Gly Ala Ser Ser Asp Thr
 2120 2125 2130
 Asn Ile Ile Lys Arg Asn Thr Lys Thr Arg Val Gly Lys Thr Thr
 2135 2140 2145
 Met Ser Asp Glu Gly Phe Gly Glu Glu Ala Glu Ile Thr Ala Asp
 2150 2155 2160
 Ser Lys Gln Gly Ile Ser Ser Phe Gly Val Gly Val Ala Ala Ala
 2165 2170 2175
 Gly Val Gly Ala Gly Val Ala Gly Thr Val Ser Val Asn Gln Phe
 2180 2185 2190
 Ala Gly Lys Thr Glu Val Asp Val Glu Glu Ala Lys Ile Leu Val
 2195 2200 2205

PCT302~1

Lys Lys Ala Glu Ile Thr Ala Lys Arg Tyr Ser Ser Val Ala Ile
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 Gly Asn Ala Ala Val Gly Val Ala Ala Lys Gly Ala Gly Ile Gly
 2225 2230 2235
 Ala Ala Val Ala Val Thr Lys Asp Glu Ser Asn Thr Arg Ala Arg
 2240 2245 2250
 Val Lys Asn Ser Lys Ile Met Thr Arg Asn Lys Leu Asp Val Ile
 2255 2260 2265
 Ala Glu Asn Glu Ile Lys Ser Gly Thr Gly Ile Gly Ser Ala Gly
 2270 2275 2280
 Ala Gly Ile Leu Ala Ala Gly Val Ser Gly Val Val Ser Val Asn
 2285 2290 2295
 Asn Ile Ala Asn Lys Val Glu Thr Asp Ile Asp His Ser Thr Leu
 2300 2305 2310
 His Ser Ser Thr Asp Val Asn Val Lys Ala Leu Asn Lys Ile Ser
 2315 2320 2325
 Asn Ser Leu Thr Ala Gly Gly Ala Ala Gly Leu Ala Ala Val
 2330 2335 2340
 Thr Gly Val Val Ser Val Asn Thr Ile Asn Ser Ser Val Ile Ala
 2345 2350 2355
 Arg Val His Asn Asn Ser Asp Leu Thr Ser Val Arg Glu Lys Val
 2360 2365 2370
 Asn Val Thr Ala Lys Glu Glu Lys Asn Ile Lys Gln Thr Ala Ala
 2375 2380 2385
 Asn Ala Gly Ile Gly Gly Ala Ala Ile Gly Ala Asn Val Leu Val
 2390 2395 2400
 Asn Asn Phe Gly Thr Ala Val Glu Asp Arg Lys Asn Ser Glu Gly
 2405 2410 2415
 Lys Gly Thr Glu Val Leu Lys Thr Leu Asp Glu Val Asn Lys Glu
 2420 2425 2430
 Gln Asp Lys Lys Val Asn Asp Ala Thr Lys Lys Ile Leu Gln Ser
 2435 2440 2445
 Ala Gly Ile Ser Thr Glu Asp Thr Ser Val Lys Ala Asp Arg Gly
 2450 2455 2460
 Asp Thr Gln Gly Glu Gly Ile Lys Ala Ile Val Lys Thr Ser Asp
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 Ile Ile Gly Lys Asn Val Asp Ile Thr Thr Glu Asp Lys Asn Asn
 2480 2485 2490
 Ile Thr Ser Thr Gly Gly Leu Gly Thr Ala Gly Leu Ala Ser Ala
 2495 2500 2505
 Ser Gly Thr Val Ala Val Thr Asn Ile Lys Arg Asn Ser Gly Val
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 Thr Val Glu Asn Ser Phe Val Lys Ala Ala Glu Lys Val Asn Val
 2525 2530 2535
 Arg Ser Asp Ile Thr Gly Asn Val Ala Leu Thr Ala Tyr Gln Gly

PCT302~1

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Pro Val Gly Ala Leu Gly Ile	Gly Ala Ala Tyr Ala	Glu Leu Asn
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Ser Asn Gly Arg Ser Asn Ile	Ser Ile Lys Asn Ser	Lys Leu Leu
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Gly Lys Asn Ile Asp Val Ile	val Lys Asp Lys Ser	Glu Leu Arg
2585 2590	2595	
Ala Glu Ala Lys Gly Leu Thr	val Gly Ala Val Ala	Ala Gly Ala
2600 2605	2610	
Ile Ile Ser Lys Ala Lys Asn	Glu Met Asn Ser Glu	Val Glu Ile
2615 2620	2625	
Glu Lys Ser Ile Phe Asn Glu	Glu Asn Arg Val Thr	Ser Pro Ser
2630 2635	2640	
Lys Gly Ile Gly Arg Glu Ile	Asn Val Lys Val Glu	Lys Glu Asn
2645 2650	2655	
Arg Val Thr Ala Glu Ser Gln	Gly Ala Ser Val Gly	Ala Val Ala
2660 2665	2670	
Gly Ala Gly Ile Ile Ser Glu	Ala Lys Asp Ala Gly	Ser Ser Tyr
2675 2680	2685	
Leu Lys Val Ser Thr Lys Ser	Gly Arg Ser Ile Phe	His Ala Asp
2690 2695	2700	
Asn Val Asn Met Glu Ala Thr	His Lys Met Lys Val	Thr Ala Val
2705 2710	2715	
Ser Lys Ala Val Thr Gly Ser	Val Leu Gly Gly Val	Gly Val Thr
2720 2725	2730	
Lys Ala Glu Ala Thr Ala Ala	Gly Lys Thr Met Val	Glu Val Glu
2735 2740	2745	
Glu Gly Asn Leu Phe Arg Thr	Asn Arg Leu Asn Ala	Ile Ser Lys
2750 2755	2760	
Val Glu Gly Leu Asp Glu Asp	Lys Val Thr Ala Lys	Ser Ser Val
2765 2770	2775	
Val Ser Gly Asn Gly Gly	Ile Ala Gly Ala Gly	Val Asn Thr
2780 2785	2790	
Ser Thr Ala Gln Ser Asn Thr	Glu Ser Val Val Arg	Leu Arg Lys
2795 2800	2805	
Gln Asp Tyr Glu Asn Asn Asp	Tyr Thr Lys Lys Tyr	Ile Ser Glu
2810 2815	2820	
Val Asn Ala Leu Ala Leu Asn	Asp Thr Lys Asn Glu	Ala Asn Ile
2825 2830	2835	
Glu Ser Leu Ala Val Ala Gly	Val His Ala Gln Gly	Thr Asn Lys
2840 2845	2850	
Ala Phe Thr Arg Ser Asn Lys	Leu Thr Ser Thr Thr	Val Asn Gly
2855 2860	2865	
Gly Asn Val Ser Gln Leu Arg	Ala Lys Ala Leu Ala	Lys Asn Glu
2870 2875	2880	

PCT302~1

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 2890 2895
 Glu Thr Ala Ala Val Glu Asn 2900 Tyr Thr Lys Ser Thr 2910 Thr Gly Ala
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 Leu Val Ala Gly Asn Trp Glu 2915 Ile Gly Asp Lys Leu Glu Thr Ile
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 Ala Arg Asp Asn Thr Ile Val 2930 Arg Val Asn Gly Asp Gly Thr Lys
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 Ser Gly Glu Thr Lys Ser Ser 2960 Ile Glu Asp Lys Ala Arg Ile Val
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 Gly Thr Gly Ser Val Asn Val 2975 Asp Ala Leu Asn Glu Leu Asp Val
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 Asp Leu Gln Gly Lys Ser Gly 2990 Tyr Gly Gly Ile Gly Ile Gly
 2995 3000
 Asn Val Asp Val Asn Asn Val 3005 Ile Lys Lys Asn Val Glu Ala Lys
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 Ile Gly Arg His Ala Ile Val 3020 Glu Thr Thr Gly Lys Gln Glu Tyr
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 Gln Ala Phe Thr Arg Ala Lys 3035 Val Asn Ile Leu Gly Lys Gly Asp
 3040 3045
 Ala Ala Ala Ala Ala Ile Ser Asn Val His Ile Ser Asn Glu
 3050 3055 3060
 Met Asp Ile Lys Asn Leu Ala 3065 Lys Gln Tyr Ala Ser Ser Gln Leu
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 3085 3090
 Glu Ser Asn Val Asn Val His 3095 Gly Val Ala Glu Ala Arg Gly Ala
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 Gly Ala Lys Ala Thr Val Ser 3110 Val Lys Asn Gln Ile Asn Arg Thr
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 Asn Asn Val Asp Leu Ala Gly 3125 Lys Ile Lys Thr Glu Gly Asn Ile
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 Asn Val Tyr Ala Gly Tyr Asp 3140 Lys Asn Tyr Asn Ile Ser Lys Thr
 3145 3150
 Asn Ser Lys Ala Ile Ala Asp 3155 Ala Lys Ser His Ala Ala Ala
 3160 3165
 Ser Ala Thr Ala Thr Ile Glu 3170 Lys Asn Glu Val Lys Phe Asn Asn
 3175 3180
 Ala Ile Arg Glu Phe Lys Asn 3185 Asn Leu Ala Arg Leu Glu Gly Lys
 3190 3195
 Ala Asn Lys Lys Thr Ser Val 3200 Gly Ser Asn Gln Val Asp Trp Tyr
 3205 3210
 Thr Asp Lys Tyr Thr Trp His 3215 Ser Ser Glu Lys Ala Tyr Lys Lys
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PCT302~1

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<210> 2

<211> 369

<212> PRT

<213> *Fusobacterium necrophorum*

<400> 2

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35 40 45

Lys Ile Glu Lys Lys Asp Asn Val Tyr Asp Ile Thr Thr Asn Lys Ile
50 55 60

Gln Gly Glu Asn Ala Phe Asn Ser Phe Asn Arg Phe Ala Leu Thr Glu
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Asn Asn Ile Ala Asn Leu Tyr Phe Gly Glu Lys Asn Ser Thr Gly Val
85 90 95

Asn Asn Leu Phe Asn Phe Val Asn Gly Lys Ile Glu Val Asp Gly Ile
100 105 110

Ile Asn Gly Ile Arg Glu Asn Lys Ile Gly Gly Asn Leu Tyr Phe Leu
115 120 125

Ser Ser Glu Gly Met Ala Val Gly Lys Asn Gly Val Ile Asn Ala Gly
130 135 140

Ser Phe His Ser Ile Ile Pro Lys Gln Asp Asp Phe Lys Lys Ala Leu
145 150 155 160

Glu Glu Ala Lys His Gly Lys Val Phe Asn Gly Ile Ile Pro Val Asp
165 170 175

Gly Lys Val Lys Ile Pro Leu Asn Pro Asn Gly Ser Ile Thr Val Glu
180 185 190

Gly Lys Ile Asn Ala Val Glu Gly Ile Gly Leu Tyr Ala Ala Asp Ile
195 200 205

Arg Leu Lys Asp Thr Ala Ile Leu Lys Thr Gly Ile Thr Asp Phe Lys
210 215 220

Asn Leu Val Asn Ile Ser Asp Arg Ile Asn Ser Gly Leu Thr Gly Asp
225 230 235 240

Leu Lys Ala Thr Lys Thr Lys Ser Gly Asp Ile Ile Leu Ser Ala His
245 250 255

Ile Asp Ser Pro Gln Lys Ala Met Gly Lys Asn Ser Thr Val Gly Lys
260 265 270

Arg Ile Glu Glu Tyr Val Lys Gly Asn Thr Lys Ala Asn Ile Glu Ser
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PCT302~1

275	280	285	
Asp Ala Val Leu Glu Ala Asp Gly Asn Ile Lys Ile Ser Ala Lys Ala			
290	295	300	
Thr Asn Gly Arg Phe Ile Lys Lys Glu Gly Glu Lys Glu Thr Tyr Asn			
305	310	315	320
Thr Pro Leu Ser Leu Ser Asp Val Glu Ala Ser Val Arg Val Asn Lys			
325	330	335	
Gly Lys Val Ile Gly Lys Asn Val Asp Ile Thr Ala Glu Ala Lys Asn			
340	345	350	
Phe Tyr Asp Ala Thr Leu Val Thr Lys Leu Ala Lys His Ser Phe Ser			
355	360	365	

Phe

<210> 3

<211> 927

<212> PRT

<213> *Fusobacterium necrophorum*

<400> 3

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20	25	30	
Val Ile Gly Lys Asn Val Asp Ile Thr Ala Glu Ala Lys Asn Phe Tyr			
35	40	45	
Asp Ala Thr Leu Val Thr Lys Leu Ala Lys His Ser Phe Ser Phe Val			
50	55	60	
Thr Gly Ser Ile Ser Pro Ile Asn Leu Asn Gly Phe Leu Gly Leu			
65	70	75	80
Thr Ser Lys Ser Ser Val Val Ile Gly Lys Asp Ala Lys Val Glu Ala			
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115	120	125	
Lys Ala Asn Gly Lys Leu Leu Ser Ile Gly Ala Gly Tyr Ile Ser Ala			
130	135	140	
Lys Ser Asn Ser Asn Val Thr Ile Glu Gly Glu Val Lys Ser Lys Gly			
145	150	155	160
Arg Ala Asp Ile Thr Ser Lys Ser Glu Asn Thr Ile Asp Ala Ser Val			
165	170	175	
Ser Val Gly Thr Met Arg Asp Ser Asn Lys Val Ala Leu Ser Val Leu			
180	185	190	

PCT302~1

Val Thr Glu Gly Glu Asn Lys Ser Ser Val Lys Ile Ala Lys Gly Ala
 195 200 205
 Lys Val Glu Ser Glu Thr Asp Asp Val Asn Val Arg Ser Glu Ala Ile
 210 215 220
 Asn Ser Ile Arg Ala Ala Val Lys Gly Gly Leu Gly Asp Ser Gly Asn
 225 230 235 240
 Gly Val Val Ala Ala Asn Ile Ser Asn Tyr Asn Ala Ser Ser Arg Ile
 245 250 255
 Asp Val Asp Gly Tyr Leu His Ala Lys Lys Arg Leu Asn Val Glu Ala
 260 265 270
 His Asn Ile Thr Lys Asn Ser Val Leu Gln Thr Gly Ser Asp Leu Gly
 275 280 285
 Thr Ser Lys Phe Met Asn Asp His Val Tyr Glu Ser Gly His Leu Lys
 290 295 300
 Ser Ile Leu Asp Ala Ile Lys Gln Arg Phe Gly Gly Asp Ser Val Asn
 305 310 315 320
 Glu Glu Ile Lys Asn Lys Leu Thr Asn Leu Phe Ser Val Gly Val Ser
 325 330 335
 Ala Thr Ile Ala Asn His Asn Asn Ser Ala Ser Val Ala Ile Gly Glu
 340 345 350
 Ser Gly Arg Leu Ser Ser Gly Val Glu Gly Ser Asn Val Arg Ala Leu
 355 360 365
 Asn Glu Ala Gln Asn Leu Arg Ala Thr Thr Ser Ser Gly Ser Val Ala
 370 375 380
 Val Arg Lys Glu Glu Lys Lys Leu Ile Gly Asn Ala Ala Val Phe
 385 390 395 400
 Tyr Gly Asn Tyr Lys Asn Asn Ala Ser Val Thr Ile Ala Asp His Ala
 405 410 415
 Glu Leu Val Ser Glu Gly Lys Ile Asp Ile Asn Ser Glu Asn Lys Ile
 420 425 430
 Glu Tyr Lys Asn Pro Ser Lys Met Ala Lys Ser Val Ile Asp Lys Leu
 435 440 445
 Glu Leu Leu Lys Arg Ala Phe Gly Lys Glu Thr Lys Thr Pro Glu Tyr
 450 455 460
 Asp Pro Lys Asp Ile Glu Ser Ile Glu Lys Leu Leu Asn Ala Phe Ser
 465 470 475 480
 Glu Lys Leu Asp Gly Lys Pro Glu Leu Leu Leu Asn Gly Glu Arg Met
 485 490 495
 Thr Ile Ile Leu Pro Asp Gly Thr Ser Lys Thr Gly Thr Ala Ile Glu
 500 505 510
 Ile Ala Asn Tyr Val Gln Gly Glu Met Lys Lys Leu Glu Glu Lys Leu
 515 520 525
 Pro Lys Gly Phe Lys Ala Phe Ser Glu Gly Leu Ser Gly Leu Ile Lys
 530 535 540
 Glu Thr Leu Asn Phe Thr Gly Val Gly Asn Tyr Ala Asn Phe His Thr
 545 550 555 560

PCT302~1

Phe Thr Ser Ser Gly Ala Asn Gly Glu Arg Asp Val Ser Ser Val Gly
565 570 575
Gly Ala Val Ser Trp Val Glu Gln Glu Asn Tyr Ser Lys Val Ser Val
580 585 590
Gly Lys Gly Ala Lys Leu Ala Ala Lys Lys Asp Leu Asn Ile Lys Ala
595 600 605
Ile Asn Lys Ala Glu Thr Val Asn Leu Val Gly Asn Ile Gly Leu Ala
610 615 620
Arg Ser Ser Thr Ser Gly Ser Ala Val Gly Gly Arg Leu Asn Val Gln
625 630 635 640
Arg Ser Lys Asn Ser Ala Ile Val Glu Ala Lys Glu Lys Ala Glu Leu
645 650 655
Ser Gly Glu Asn Ile Asn Ala Asp Ala Leu Asn Arg Leu Phe His Val
660 665 670
Ala Gly Ser Phe Asn Gly Gly Ser Gly Gly Asn Ala Ile Asn Gly Met
675 680 685
Gly Ser Tyr Ser Gly Gly Ile Ser Lys Ala Arg Val Ser Ile Asp Asp
690 695 700
Glu Ala Tyr Leu Lys Ala Asn Lys Lys Ile Ala Leu Asn Ser Lys Asn
705 710 715 720
Asp Thr Ser Val Trp Asn Ala Ala Gly Ser Ala Gly Ile Gly Thr Lys
725 730 735
Asn Ala Ala Val Gly Val Ala Val Ala Val Asn Asp Tyr Asp Ile Ser
740 745 750
Asn Lys Ala Ser Ile Glu Asp Asn Asp Glu Gly Gln Ser Lys Tyr Asp
755 760 765
Lys Asn Lys Asp Asp Glu Val Thr Val Thr Ala Glu Ser Leu Glu Val
770 775 780
Asp Ala Lys Thr Thr Gly Thr Ile Asn Ser Ile Ser Val Ala Gly Gly
785 790 795 800
Ile Asn Lys Val Gly Ser Lys Pro Ser Glu Glu Lys Pro Lys Ser Glu
805 810 815
Glu Arg Pro Glu Gly Phe Phe Gly Lys Ile Gly Asn Lys Val Asp Ser
820 825 830
Val Lys Asn Lys Ile Thr Asp Ser Met Asp Ser Leu Thr Glu Lys Ile
835 840 845
Thr Asn Tyr Ile Ser Glu Gly Val Lys Lys Ala Gly Asn Leu Pro Ser
850 855 860
Asn Val Ser His Thr Pro Asp Lys Gly Pro Ser Phe Ser Leu Gly Ala
865 870 875 880
Ser Gly Ser Val Ser Phe Asn Asn Ile Lys Lys Glu Thr Ser Ala Val
885 890 895
Val Asp Gly Val Lys Ile Asn Leu Lys Gly Ala Asn Lys Lys Val Glu
900 905 910
Val Thr Ser Ser Asp Ser Thr Phe Val Gly Ala Trp Gly Ser

915

920

925

<210> 4

<211> 580

<212> PRT

<213> **Fusobacterium necrophorum**

<400> 4

Gly Ala Ser Gly Ser Val Ser Asn Asn Lys Lys Thr Ser Ala Val Val
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Asp Gly Val Lys Asn Lys Gly Ala Asn Lys Lys Val Val Thr Ser Ser
 20 25 30

Asp Ser Thr Val Gly Ala Trp Gly Gly Ser Ala Ala Trp Asn His Gly
 35 40 45

Ser Gly Asn Ser Asn Ser Ala Gly Ala Gly Ala Ala Val Asn Asn
 50 55 60

Ser Lys Thr Ser Ala Val Lys Asn Ser Asp Arg Asn Ala Asn Lys Lys
 65 70 75 80

Val Asn Ala Ser Gly Gly Thr Val Ala Ala Gly Ala Gly Ala Val Lys
 85 90 95

Ser Gly Gly Lys Ser Tyr Gly Thr Ser Ala Ser Asn Val Asn Asn
 100 105 110

Val Ser Ala Lys Ser Asn Asn Thr Val Ala Gly Ser Ser Lys Met Asp
 115 120 125

Val Asp Val Thr Ala Tyr Ala Asp Thr Val Thr Gly Ala Asn Ala Gly
 130 135 140

Lys Ser Asn Gly Thr Val Gly Ala Thr Val Thr Val Ala Lys Asn Asn
 145 150 155 160

Lys Val Asn Ala Ser Ser Gly Gly Arg Tyr Thr Asn Val Asn Arg Ala
 165 170 175

Asp Ala Lys Ala Ala Thr Thr Val Thr Ala Ala Val Thr Thr Gly Gly
 180 185 190

Thr Ser Ser Gly Ala Gly Gly Asn Tyr Gly Ala Val Ser Val Asn Lys
 195 200 205

Asp Asn Asp Val Ala Ser Val Asp Lys Ser Ser Gly Ala Asn Asn Val
 210 215 220

Ala Lys Asp Val Lys Gly Ser Ser Asp Ala Lys Tyr Ala Asn Gly Lys
 225 230 235 240

Asp Lys Lys Tyr Asp Arg Gly Asn Thr Thr Gly Asn Gly Tyr Tyr Thr
 245 250 255

Lys Lys Ala Lys Lys Lys Gly Ala Val Val Asn Ala Ala Ser Val Ala
 260 265 270

Gly Thr Asp Lys Ser Ala Gly Gly Val Ala Ala Val Asn Thr Val Lys
 275 280 285

PCT302~1

Asn Lys Lys Ala Ser Gly Ser Asn Lys Ala Gly Asp Lys His Ala Lys
 290 295 300

His Val Asn Val Ala Lys Ser Ser Thr Val Val Val Asn Ala Ala Ser
 305 310 315 320

Gly Ala Ser Lys Asp Ala Ser Gly Met Gly Ser Gly Ala Trp Asp Ser
 325 330 335

Asn Asp Thr Ala Lys Val Asp Lys Gly Arg Ser Ala Asp Ser Asn Val
 340 345 350

Asn Ala Asn Asn Ser Gly Val Asn Val Ala Gly Thr Ala Gly Ser Ser
 355 360 365

Thr Ala Val Gly Ala Ala Ala Asn Asn Thr His Asn Lys Thr Ser Ala
 370 375 380

Thr Gly Thr Lys Val Asn Ser Gly Lys Asn Thr Lys Val Asn Val Ala
 385 390 395 400

Asn Asp Ser His Thr Asn Val Ser Ala Gly Gly Ala Ala Ser Lys Ala
 405 410 415

Gly Gly Gly Met Val Ser Val Asn Arg Gly Ser Asp Thr Ala Val Ser
 420 425 430

Asp Ser Gly Val Ser Ser Asn Val Asp Ala Lys Asp Lys Thr Asn Thr
 435 440 445

Ala Gly Asn Ala Asn Gly Gly Lys Ala Ala Gly Val Gly Ala Thr Val
 450 455 460

Ala His Thr Asn Gly Lys Ser Val Ala Val Lys Asn Ser Lys Thr Thr
 465 470 475 480

Ala Asn Asp Asp Arg Lys Asn Asn Val Thr Ala Lys Asp Tyr Thr Met
 485 490 495

Thr Asn Thr Ala Val Gly Val Gly Ala Lys Gly Ala Ser Val Gly
 500 505 510

Ala Ser Ala Ser Thr Thr Asn Lys Thr Val Ser Ser His Val Asp Thr
 515 520 525

Asp Asp Lys Asp Asn Asn Gly Asn Lys Lys Ala Asn Val Asn Val Ala
 530 535 540

Asn Thr Ser Val Val Thr Asn Ala Thr Val Ser Gly Ala Ser Gly Ala
 545 550 555 560

Ala Val Gly Ala Gly Val Ala Val Asn Lys Thr Asn Thr Ser Ala His
 565 570 575

Lys Asn Ser Thr
 580

<210> 5

<211> 628

<212> PRT

<213> *Fusobacterium necrophorum*

<400> 5

PCT302~1

Ala val Gly Ala Gly val Ala val Asn Lys Ile Thr Gln Asn Thr Ser
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Ala His Ile Lys Asn Ser Thr Gln Asn Val Arg Asn Ala Leu Val Lys
20 25 30

Ser Lys Ser His Ser Ser Ile Lys Thr Ile Gly Ile Gly Ala Gly val
35 40 45

Gly Ala Gly Gly Ala Gly Val Thr Gly Ser Val Ala Val Asn Lys Ile
50 55 60

val Asn Asn Thr Ile Ala Glu Leu Asn His Ala Lys Ile Thr Ala Lys
65 70 75 80

Gly Asn Val Gly Val Ile Thr Glu Ser Asp Ala Val Ile Ala Asn Tyr
85 90 95

Ala Gly Thr Val Ser Gly Val Ala Arg Ala Ala Ile Gly Ala Ser Thr
100 105 110

Ser Val Asn Glu Ile Thr Gly Ser Thr Lys Ala Tyr Val Lys Asp Ser
115 120 125

Thr Val Ile Ala Lys Glu Glu Thr Asp Asp Tyr Ile Thr Thr Gln Gly
130 135 140

Gln Val Asp Lys Val Val Asp Lys Val Phe Lys Asn Leu Asn Ile Asn
145 150 155 160

Glu Asp Leu Ser Gln Lys Arg Lys Ile Ser Asn Lys Lys Gly Phe Val
165 170 175

Thr Asn Ser Ser Ala Thr His Thr Leu Lys Ser Leu Leu Ala Asn Ala
180 185 190

Ala Gly Ser Gly Gln Ala Gly Val Ala Gly Thr Val Asn Ile Asn Lys
195 200 205

Val Tyr Gly Glu Thr Glu Ala Leu Val Glu Asn Ser Ile Leu Asn Ala
210 215 220

Lys His Tyr Ser Val Lys Ser Gly Asp Tyr Thr Asn Ser Ile Gly val
225 230 235 240

Val Gly Ser Val Gly Val Gly Asn Val Gly Val Gly Ala Ser Ser
245 250 255

Asp Thr Asn Ile Ile Lys Arg Asn Thr Lys Thr Arg Val Gly Lys Thr
260 265 270

Thr Met Ser Asp Glu Gly Phe Gly Glu Glu Ala Glu Ile Thr Ala Asp
275 280 285

Ser Lys Gln Gly Ile Ser Ser Phe Gly Val Gly Val Ala Ala Ala Gly
290 295 300

Val Gly Ala Gly Val Ala Gly Thr Val Ser Val Asn Gln Phe Ala Gly
305 310 315 320

Lys Thr Glu Val Asp Val Glu Glu Ala Lys Ile Leu Val Lys Lys Ala
325 330 335

Glu Ile Thr Ala Lys Arg Tyr Ser Ser Val Ala Ile Gly Asn Ala Ala
340 345 350

Val Gly Val Ala Ala Lys Gly Ala Gly Ile Gly Ala Ala Val Ala Val

PCT302~1

355

360

365

Thr Lys Asp Glu Ser Asn Thr Arg Ala Arg Val Lys Asn Ser Lys Ile
 370 375 380

Met Thr Arg Asn Lys Leu Asp Val Ile Ala Glu Asn Glu Ile Lys Ser
 385 390 395 400

Gly Thr Gly Ile Gly Ser Ala Gly Ala Gly Ile Leu Ala Ala Gly Val
 405 410 415

Ser Gly Val Val Ser Val Asn Asn Ile Ala Asn Lys Val Glu Thr Asp
 420 425 430

Ile Asp His Ser Thr Leu His Ser Ser Thr Asp Val Asn Val Lys Ala
 435 440 445

Leu Asn Lys Ile Ser Asn Ser Leu Thr Ala Gly Gly Ala Ala Gly
 450 455 460

Leu Ala Ala Val Thr Gly Val Val Ser Val Asn Thr Ile Asn Ser Ser
 465 470 475 480

Val Ile Ala Arg Val His Asn Asn Ser Asp Leu Thr Ser Val Arg Glu
 485 490 495

Lys Val Asn Val Thr Ala Lys Glu Glu Lys Asn Ile Lys Gln Thr Ala
 500 505 510

Ala Asn Ala Gly Ile Gly Gly Ala Ala Ile Gly Ala Asn Val Leu Val
 515 520 525

Asn Asn Phe Gly Thr Ala Val Glu Asp Arg Lys Asn Ser Glu Gly Lys
 530 535 540

Gly Thr Glu Val Leu Lys Thr Leu Asp Glu Val Asn Lys Glu Gln Asp
 545 550 555 560

Lys Lys Val Asn Asp Ala Thr Lys Lys Ile Leu Gln Ser Ala Gly Ile
 565 570 575

Ser Thr Glu Asp Thr Ser Val Lys Ala Asp Arg Gly Asp Thr Gln Gly
 580 585 590

Glu Gly Ile Lys Ala Ile Val Lys Thr Ser Asp Ile Ile Gly Lys Asn
 595 600 605

Val Asp Ile Thr Thr Glu Asp Lys Asn Asn Ile Thr Ser Thr Gly Gly
 610 615 620

Leu Gly Thr Ala
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<210> 6

<211> 773

<212> PRT

<213> *Fusobacterium necrophorum*

<400> 6

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PCT302~1

Asp Ile Thr Thr Glu Asp Lys Asn Asn Ile Thr Ser Thr Gly Gly Leu
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Gly Thr Ala Gly Leu Ala Ser Ala Ser Gly Thr Val Ala Val Thr Asn
 35 40 45

Ile Lys Arg Asn Ser Gly Val Thr Val Glu Asn Ser Phe Val Lys Ala
 50 55 60

Ala Glu Lys Val Asn Val Arg Ser Asp Ile Thr Gly Asn Val Ala Leu
 65 70 75 80

Thr Ala Tyr Gln Gly Pro Val Gly Ala Leu Gly Ile Gly Ala Ala Tyr
 85 90 95

Ala Glu Leu Asn Ser Asn Gly Arg Ser Asn Ile Ser Ile Lys Asn Ser
 100 105 110

Lys Leu Leu Gly Lys Asn Ile Asp Val Ile Val Lys Asp Lys Ser Glu
 115 120 125

Leu Arg Ala Glu Ala Lys Gly Leu Thr Val Gly Ala Val Ala Ala Gly
 130 135 140

Ala Ile Ile Ser Lys Ala Lys Asn Glu Met Asn Ser Glu Val Glu Ile
 145 150 155 160

Glu Lys Ser Ile Phe Asn Glu Glu Asn Arg Val Thr Ser Pro Ser Lys
 165 170 175

Gly Ile Gly Arg Glu Ile Asn Val Lys Val Glu Lys Glu Asn Arg Val
 180 185 190

Thr Ala Glu Ser Gln Gly Ala Ser Val Gly Ala Val Ala Gly Ala Gly
 195 200 205

Ile Ile Ser Glu Ala Lys Asp Ala Gly Ser Ser Tyr Leu Lys Val Ser
 210 215 220

Thr Lys Ser Gly Arg Ser Ile Phe His Ala Asp Asn Val Asn Met Glu
 225 230 235 240

Ala Thr His Lys Met Lys Val Thr Ala Val Ser Lys Ala Val Thr Gly
 245 250 255

Ser Val Leu Gly Gly Val Gly Val Thr Lys Ala Glu Ala Thr Ala Ala
 260 265 270

Gly Lys Thr Met Val Glu Val Glu Glu Gly Asn Leu Phe Arg Thr Asn
 275 280 285

Arg Leu Asn Ala Ile Ser Lys Val Glu Gly Leu Asp Glu Asp Lys Val
 290 295 300

Thr Ala Lys Ser Ser Val Val Ser Gly Asn Gly Gly Gly Ile Ala Gly
 305 310 315 320

Ala Gly Val Asn Thr Ser Thr Ala Gln Ser Asn Thr Glu Ser Val Val
 325 330 335

Arg Leu Arg Lys Gln Asp Tyr Glu Asn Asn Asp Tyr Thr Lys Lys Tyr
 340 345 350

Ile Ser Glu Val Asn Ala Leu Ala Leu Asn Asp Thr Lys Asn Glu Ala
 355 360 365

Asn Ile Glu Ser Leu Ala Val Ala Gly Val His Ala Gln Gly Thr Asn
 370 375 380

PCT302~1

Lys Ala Phe Thr Arg Ser Asn Lys Leu Thr Ser Thr Thr Val Asn Gly
385 390 395 400
Gly Asn Val Ser Gln Leu Arg Ala Lys Ala Leu Ala Lys Asn Glu Asn
405 410 415
Tyr Gly Asn Val Lys Gly Thr Gly Ala Leu Val Gly Ala Glu Thr
420 425 430
Ala Ala Val Glu Asn Tyr Thr Lys Ser Thr Thr Gly Ala Leu Val Ala
435 440 445
Gly Asn Trp Glu Ile Gly Asp Lys Leu Glu Thr Ile Ala Arg Asp Asn
450 455 460
Thr Ile Val Arg Val Asn Gly Asp Gly Thr Lys Gly Gly Leu Val Gly
465 470 475 480
Lys Asn Gly Ile Ser Val Lys Asn Thr Ile Ser Gly Glu Thr Lys Ser
485 490 495
Ser Ile Glu Asp Lys Ala Arg Ile Val Gly Thr Gly Ser Val Asn Val
500 505 510
Asp Ala Leu Asn Glu Leu Asp Val Asp Leu Gln Gly Lys Ser Gly Gly
515 520 525
Tyr Gly Gly Ile Gly Ile Gly Asn Val Asp Val Asn Asn Val Ile Lys
530 535 540
Lys Asn Val Glu Ala Lys Ile Gly Arg His Ala Ile Val Glu Thr Thr
545 550 555 560
Gly Lys Gln Glu Tyr Gln Ala Phe Thr Arg Ala Lys Val Asn Ile Leu
565 570 575
Gly Lys Gly Asp Ala Ala Ala Ala Ala Ile Ser Asn Val His Ile
580 585 590
Ser Asn Glu Met Asp Ile Lys Asn Leu Ala Lys Gln Tyr Ala Ser Ser
595 600 605
Gln Leu Ile Thr Lys Asn Ser Lys Asn Asn Ile Thr Leu Ala Ser Ser
610 615 620
Ser Glu Ser Asn Val Asn Val His Gly Val Ala Glu Ala Arg Gly Ala
625 630 635 640
Gly Ala Lys Ala Thr Val Ser Val Lys Asn Gln Ile Asn Arg Thr Asn
645 650 655
Asn Val Asp Leu Ala Gly Lys Ile Lys Thr Glu Gly Asn Ile Asn Val
660 665 670
Tyr Ala Gly Tyr Asp Lys Asn Tyr Asn Ile Ser Lys Thr Asn Ser Lys
675 680 685
Ala Ile Ala Asp Ala Lys Ser His Ala Ala Ala Ala Ser Ala Thr Ala
690 695 700
Thr Ile Glu Lys Asn Glu Val Lys Phe Asn Asn Ala Ile Arg Glu Phe
705 710 715 720
Lys Asn Asn Leu Ala Arg Leu Glu Gly Lys Ala Asn Lys Lys Thr Ser
725 730 735
Val Gly Ser Asn Gln Val Asp Trp Tyr Thr Asp Lys Tyr Thr Trp His

740 745 PCT302~1 750

Ser Ser Glu Lys Ala Tyr Lys Lys Leu Thr Tyr Gln Ser Lys Arg Gly
755 760 765

Glu Lys Gly Lys Lys
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<210> 7

<211> 338

<212> PRT

<213> *Fusobacterium necrophorum*

<400> 7

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Gln Ile Tyr Glu Pro Lys Arg Gln Gln Phe Phe Val Phe Ala Asp Asn
20 25 30

Leu Gly Gln Lys Asn Thr Gly Glu Leu Arg Trp Gly Leu Asn Tyr Ile
35 40 45

Asn Asn Ser Val Thr Gly Asn Arg Asp Gln Leu Ser Leu Thr Ser Leu
50 55 60

Val Thr Glu Gly Thr Ala Ser Leu Ser Ser Phe Tyr Thr Phe Pro Val
65 70 75 80

Ser Lys Lys Gly Thr Lys Ile Ser Leu Gln His Ser Val Gly Lys Leu
85 90 95

Lys His Ile Gln Gly Ala Leu Lys His Lys Ile Thr Gly Asn Ser Tyr
100 105 110

Ser Tyr Gly Val Gly Ile Val His Pro Ile Leu Val His Glu Lys Asn
115 120 125

Lys Val Glu Leu Ser Leu Asp Trp Val Lys Gln Arg Thr Val Thr Asp
130 135 140

Leu Leu Lys Leu Lys Trp Val Asn Asn Arg Leu Ser Lys Tyr Thr Ala
145 150 155 160

Gly Ile Gly Ile Ser His Tyr Glu Glu Asp Ser Val Phe Tyr Thr Lys
165 170 175

Gln Asn Ile Thr Lys Gly Lys Phe Ile Pro Ile Ser Gly Asp Ala Arg
180 185 190

Asn Tyr Thr Lys Tyr Asp Met Phe Leu Ile Tyr Gln Lys Asn Leu Lys
195 200 205

Tyr Asn Thr Leu Val Thr Leu Lys Met Ala Gly Gln Tyr Ser Leu Ser
210 215 220

Lys Lys Leu Pro Ser Val Glu Gln Ile Tyr Ala Gly Gly Ala Tyr Asn
225 230 235 240

Val Arg Gly Tyr Pro Glu Asn Phe Met Gly Ala Glu His Gly Val Phe
245 250 255

<210> 8
<211> 9726
<212> DNA
<213> ***Fusobacterium necrophorum***

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accgcgactg agaattttgg aacaaaaata gaaaaaaagg ataatgttt a tacattact 180
acaaacaaga ttcaagggga gaacgcttt aacagttta atagattgc tttaacagaa 240
aataatatag caaatctata ttttgggaa aagaatagta cggggtaaa taatctttt 300
aactttgtca atggaaaaat tgaagtagat gggattatca acggaattcg agaaaataaa 360
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acagattttt aaaaatttagt caatatttagt gatcgaataa attctggctc gaccggagat 720
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PCT302~1

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PCT302~1

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PCT302~1

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PCT302~1

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PCT302~1

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<210> 9

<211> 1130

<212> DNA

<213> *Fusobacterium necrophorum*

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<210> 10

PCT302~1

<211> 2780

<212> DNA

<213> *Fusobacterium necrophorum*

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tttagcttg ttacaggttc tatttctcct atcaatttaa atggattttt aggtttattg	240
acaagtaagt ccagtgtcgt tattggaaaa gatgccaaag tcgaagcaac agaaggaaag	300
gcaaataattc attcttacag tggagtaaga gcaactatgg gagcagctac ttctccatta	360
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tccgtcaaga ttgctaaagg agcaaaagta gaatcagaaa cggatgatgt aaatgtgaga	660
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PCT302~1

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<210> 11

<211> 2141

<212> DNA

<213> *Fusobacterium necrophorum*

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aatgcttcta ttagtggtgg gagatataact aacgttaatc gagcggacgc当地 aaaagcttctt	660

PCT302~1

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<210> 12

<211> 1887

<212> DNA

<213> *Fusobacterium necrophorum*

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aaacaattgg	aattggagct	ggagttggag	ctggaggagc	tggagtgaca	ggttctgttag		180

PCT302~1

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<210> 13

<211> 2322

<212> DNA

<213> *Fusobacterium necrophorum*

PCT302~1

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